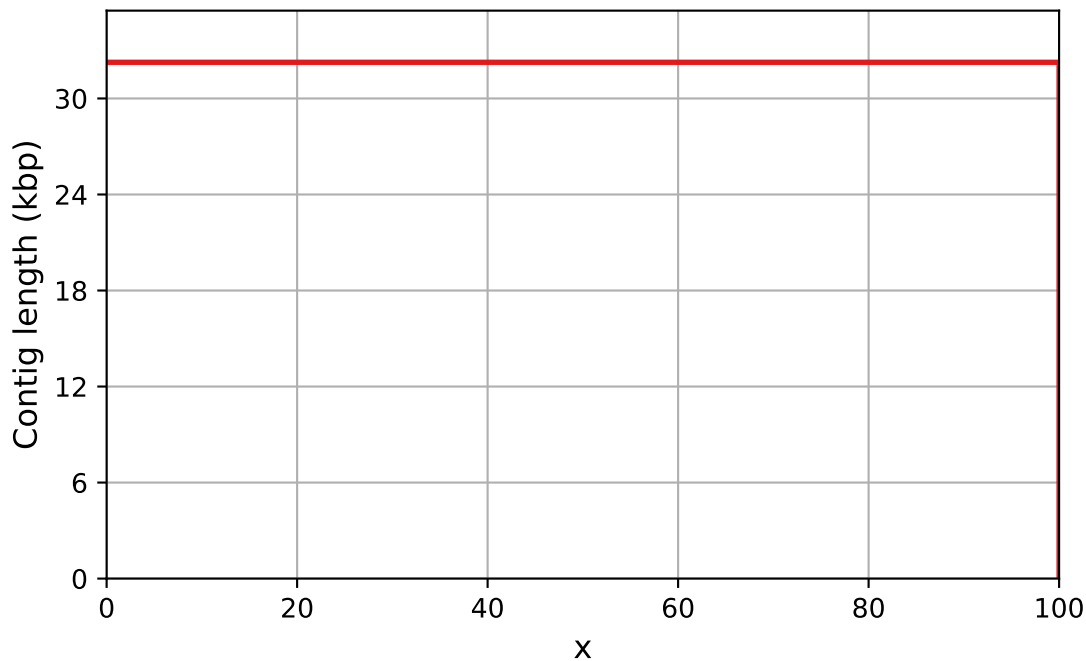


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1
# contigs (>= 1000 bp)	1
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	32261
Total length (>= 1000 bp)	32261
Total length (>= 5000 bp)	32261
Total length (>= 10000 bp)	32261
Total length (>= 25000 bp)	32261
Total length (>= 50000 bp)	0
# contigs	1
Largest contig	32261
Total length	32261
GC (%)	35.32
N50	32261
N90	32261
auN	32261.0
L50	1
L90	1
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

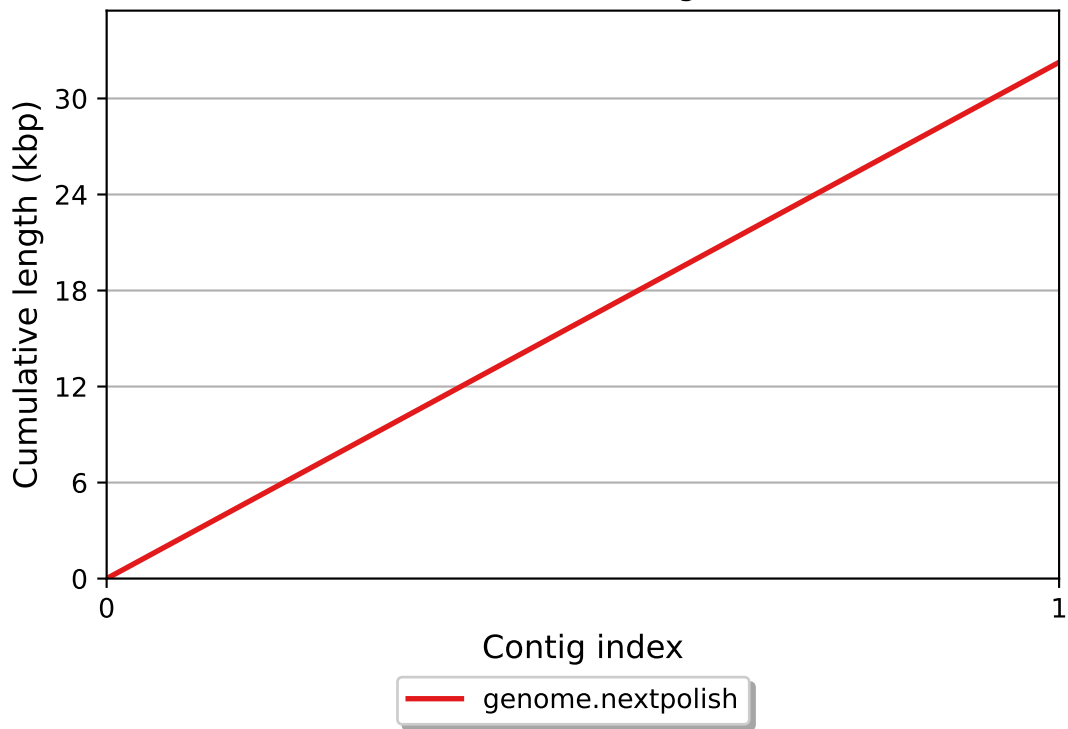
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

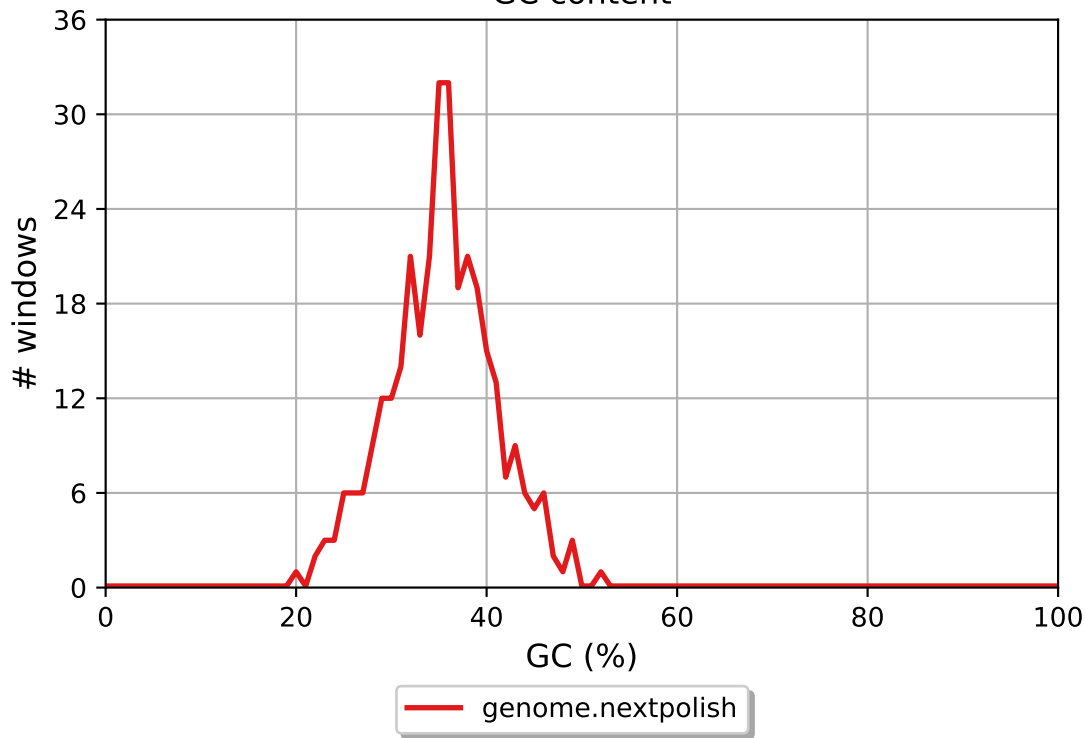


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

